

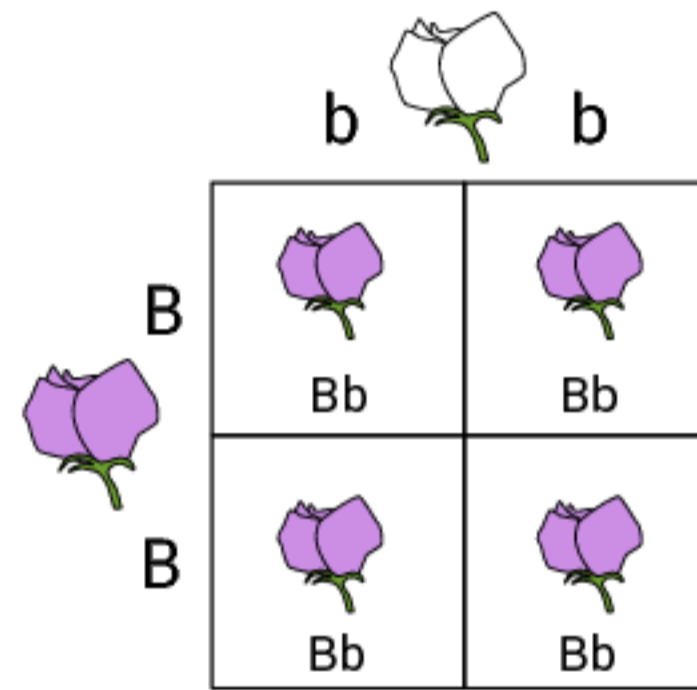
# BIOS1100 H17 uke 7

## Lex Nederbragt

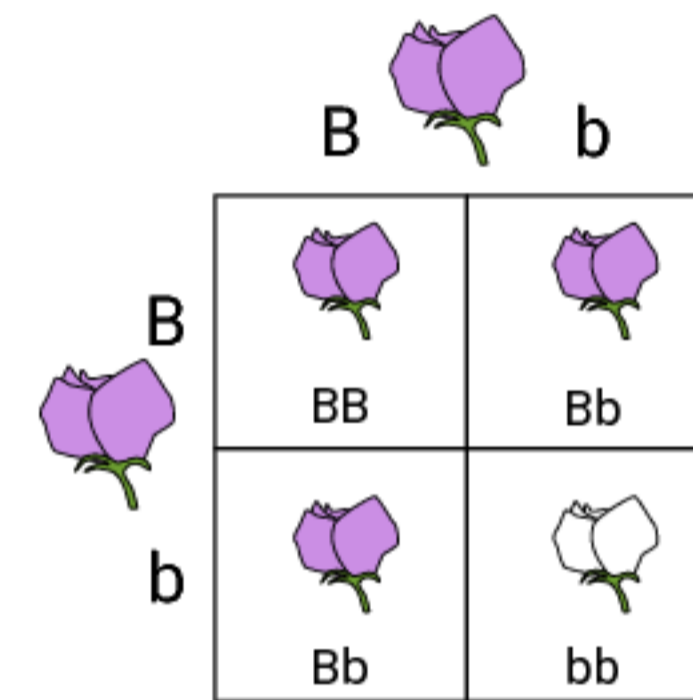
P generation



F<sub>1</sub> generation



F<sub>2</sub> generation



# Ukens forelesning

---

- noen praktiske ting
- nytt stoff denne uken
- utvalgte øvelser

# Noen praktiske ting

---

## Uke 41 (kursuke 8)

- undervisningsfri uke
- ingen forelesning
- utvidede snublegrupper
  - tirsdag 10. oktober kl 10:15-14:00 seminarrom 3508
  - torsdag 12. oktober kl 10:15-14:00 seminarrom 4619

# Noen praktiske ting

---

## Obliger:

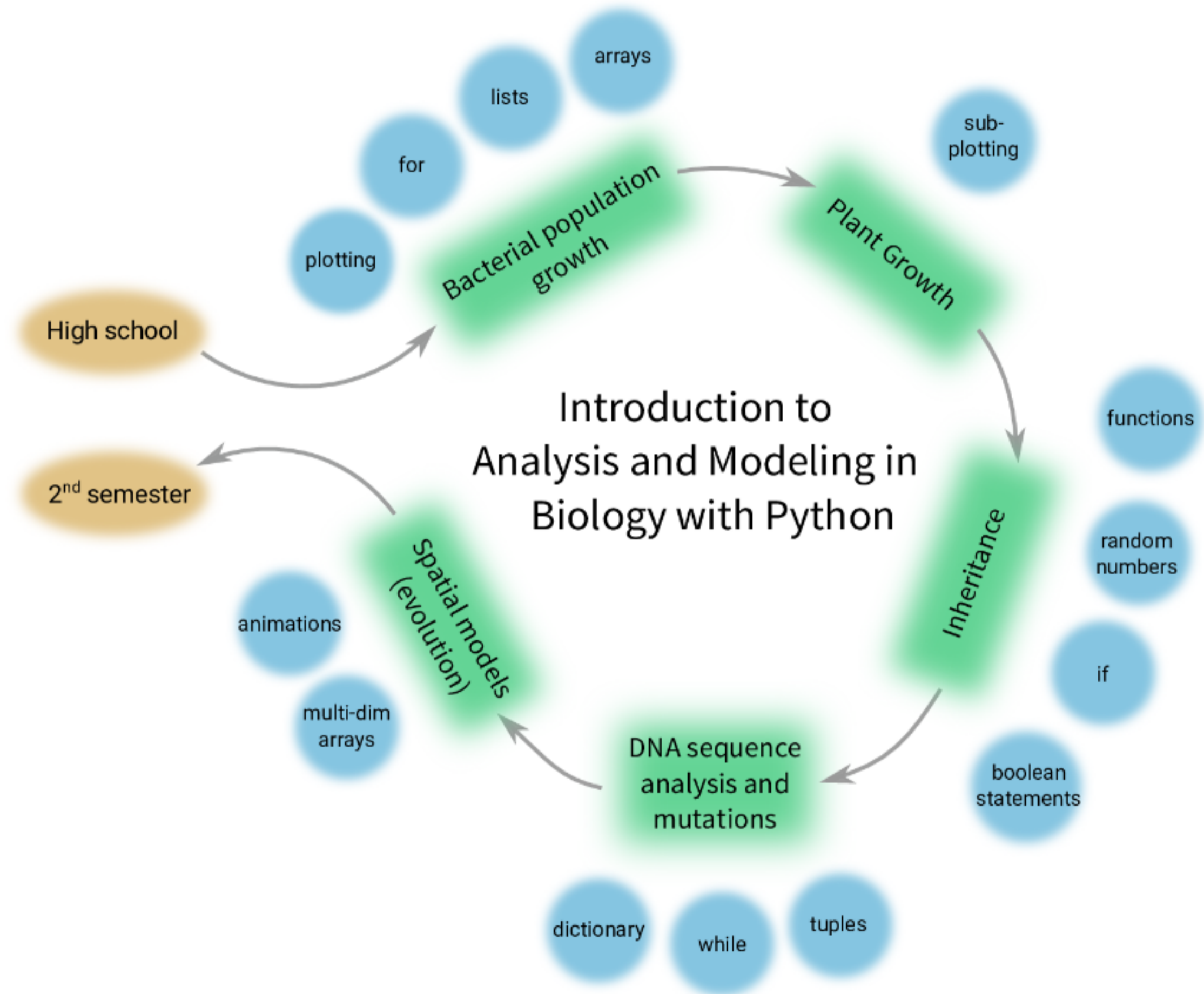
- tirsdag 3. oktober 23:59: frist oblig for uke **6** (kalenderuke 39)
- tirsdag 10. oktober 23:59: frist oblig for uke **7** (uke 40, denne uken)
- *ingen* oblig for uke **8** (kalenderuke 41)
- tirsdag 24. oktober 23:59: frist oblig for uke **9** (kalenderuke 42)

# Noen praktiske ting

---

- Turtle konkurranse!
  - send din turtle tegning som notebook til [bios1100@ibv.uio.no](mailto:bios1100@ibv.uio.no)
  - den mest kreative får en pris!
  - (gruppe)lærere er juryen
  - kodekvalitet er ikke viktig

# Undervisningsplan



# Læringsmål denne uke

---

## Biology

- kjenne antagelser for Mendels arvelover
- kunne lage Punnett diagrammer for hånd
- kunne sannsynlighetsteorien for tilfeldig parring
- kunne modellere tilfeldig parring

## Programmering

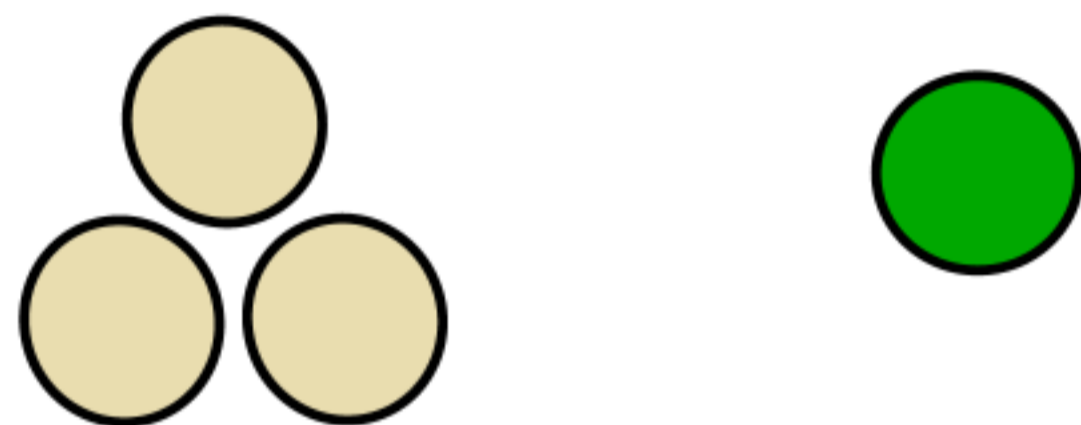
- tilfeldig valg med `choice`
- funksjoner
- `if` tester og 'boolean statements'

# Ratio

---

$\left(\frac{3}{4}\right)$  versus  $\left(\frac{1}{4}\right)$

Seeds



3 : 1



# Kombinere lister

---

```
cells = ["Skin", "Muscle", "Nerve", "Blood"]
components = ["Nucleus", "Ribosome", "Golgi apparatus", "Lysosome", "Vacuole", "Mitochondrion", "Chloroplast", "Peroxisome"]
for cell in cells:
    for component in components:
        print(cell, component)
    print("Done with components for", cell)
print("Done with all cells")
```

Skin Nucleus  
Skin Ribosome  
Skin Golgi apparatus  
Skin Lysosome  
Skin Vacuole  
Skin Mitochondrion  
Skin Chloroplast  
Skin Peroxisome  
Done with components for Skin  
Muscle Nucleus  
Muscle Ribosome  
Muscle Golgi apparatus  
Muscle Lysosome  
Muscle Vacuole  
Muscle Mitochondrion  
Muscle Chloroplast  
Muscle Peroxisome  
Done with components for Muscle  
Nerve Nucleus  
Nerve Ribosome  
Nerve Golgi apparatus  
Nerve Lysosome  
Nerve Vacuole  
Nerve Mitochondrion  
Nerve Chloroplast  
Nerve Peroxisome  
Done with components for Nerve  
Blood Nucleus  
Blood Ribosome  
Blood Golgi apparatus  
Blood Lysosome  
Blood Vacuole  
Blood Mitochondrion  
Blood Chloroplast  
Blood Peroxisome  
Done with components for Blood  
Done with all cells

## Kombinere lister

---

```
cells = ["Skin", "Muscle", "Nerve", "Blood"]
components = ["Nucleus", "Ribosome", "Golgi apparatus", "Lysosome", "Vacuole",
              "Mitochondrion", "Chloroplast", "Peroxisome"]
```

## Kombinere lister

---

```
cells = ["Skin", "Muscle", "Nerve", "Blood"]
components = ["Nucleus", "Ribosome", "Golgi apparatus", "Lysosome", "Vacuole",
"Mitochondrion", "Chloroplast", "Peroxisome"]

for cell in cells:
    for component in components:
        print(cell, component)
    print("Done with components for", cell)
print("Done with all cells")
```

## Kombinere lister

---

```
pure_violet_flower = ["B", "B"]
pure_white_flower = ["b", "b"]

print("F1 generation:")

for allele_1 in pure_violet_flower:
    for allele_2 in pure_white_flower:
        print("Possible offspring:", allele_1, allele_2)
```

```
F1 generation:
Possible offspring: B b
Possible offspring: B b
Possible offspring: B b
Possible offspring: B b
```

## Kombinere lister

---

```
crossbred_flower = ["B", "b"]

print("F2 generation:")

for allele_1 in crossbred_flower:
    for allele_2 in crossbred_flower:
        print("Possible offspring:", allele_1, allele_2)
```

```
F2 generation:
Possible offspring: B B
Possible offspring: B b
Possible offspring: b B
Possible offspring: b b
```

# Pandas dataframe

---

Columns

	colA	colB	colC
rowD			
rowE			
rowF			

# Pandas dataframe

---

```
col_names = ["colA", "colB", "colC"]  
row_names = ["rowD", "rowE", "rowF"]  
df_example = pandas.DataFrame(index=row_names, columns=col_names)
```

		Columns		
		colA	colB	colC
Indexes (rows)	rowD			
	rowE			
	rowF			

## Pandas dataframe

```
col_names = ["colA", "colB", "colC"]
row_names = ["rowD", "rowE", "rowF"]
df_example = pandas.DataFrame(index=row_names, columns=col_names)
print(df_example)
```

```
      colA colB colC
rowD  NaN  NaN  NaN
rowE  NaN  NaN  NaN
rowF  NaN  NaN  NaN
```

**Columns**

	colA	colB	colC
rowD			
rowE			
rowF			



# Pandas dataframe

---

		Columns		
		0	1	2
Rows	0	[0, 0]	[0, 1]	[0, 2]
	1	[1, 0]	[1, 1]	[1, 2]
	2	[2, 0]	[2, 1]	[2, 2]

Row-by-column indexing

# Pandas dataframe

---

Fylle tabellen:

```
iloc[row, column]
```

		Columns		
		0	1	2
Rows	0	[0, 0]	[0, 1]	[0, 2]
	1	[1, 0]	[1, 1]	[1, 2]
	2	[2, 0]	[2, 1]	[2, 2]

# Pandas dataframe

---

For eksempel:

```
df_example.iloc[0, 0] = "firstcell"  
print(df_example)
```

	colA	colB	colC
rowD	firstcell	NaN	NaN
rowE	NaN	NaN	NaN
rowF	NaN	NaN	NaN

**Columns**

	0	1	2
0	[0, 0]	[0, 1]	[0, 2]
1	[1, 0]	[1, 1]	[1, 2]
2	[2, 0]	[2, 1]	[2, 2]

**Rows**

# Python enumerate funksjonen

---

En vanlig for løkke:

```
my_list = ["A", "B", "C", "D"]
for element in my_list:
    print("element er", element)
```

```
element er A
element er B
element er C
element er D
```

# Python enumerate funksjonen

---

Men hvordan å få denne outputen?

```
index 0 er element A  
index 1 er element B  
index 2 er element C  
index 3 er element D
```

# Python enumerate funksjonen

---

Men hvordan å få denne outputen?

```
index 0 er element A
index 1 er element B
index 2 er element C
index 3 er element D
```

Mulighet 1:

```
my_list = ["A", "B", "C", "D"]

for index in range(4):
    print("index", index, "er element", my_list[index])
```

# Python enumerate funksjonen

---

Men hvordan å få denne outputen?

```
index 0 er element A
index 1 er element B
index 2 er element C
index 3 er element D
```

Mulighet 2, mer generelt:

```
my_list = ["A", "B", "C", "D"]

for index in range(len(my_list)):
    print("index", index, "er element", my_list[index])
```

# Python enumerate funksjonen

---

Men hvordan å få denne outputen?

```
index 0 er element A
index 1 er element B
index 2 er element C
index 3 er element D
```

Mulighet 3, med `enumerate`

```
my_list = ["A", "B", "C", "D"]

for index, element in enumerate(my_list):
    print("index", index, "er element", element)
```



## Python enumerate funksjonen

---

```
my_list = ["A", "B", "C", "D"]

for index, element in enumerate(my_list):
    print("index:", index, ", element:", element)
```

```
index 0 er element A
index 1 er element B
index 2 er element C
index 3 er element D
```

# Pandas dataframe

---

```
col_names = ["colA", "colB", "colC"]
row_names = ["rowD", "rowE", "rowF"]

for col, col_name in enumerate(col_names):
    for row, row_name in enumerate(row_names):
        print(col, col_name, row, row_name, col_name+row_name)
```

```
0 colA 0 rowD colArowD
0 colA 1 rowE colArowE
0 colA 2 rowF colArowF
1 colB 0 rowD colBrowD
1 colB 1 rowE colBrowE
1 colB 2 rowF colBrowF
2 colC 0 rowD colCrowD
2 colC 1 rowE colCrowE
2 colC 2 rowF colCrowF
```

# Pandas dataframe

---

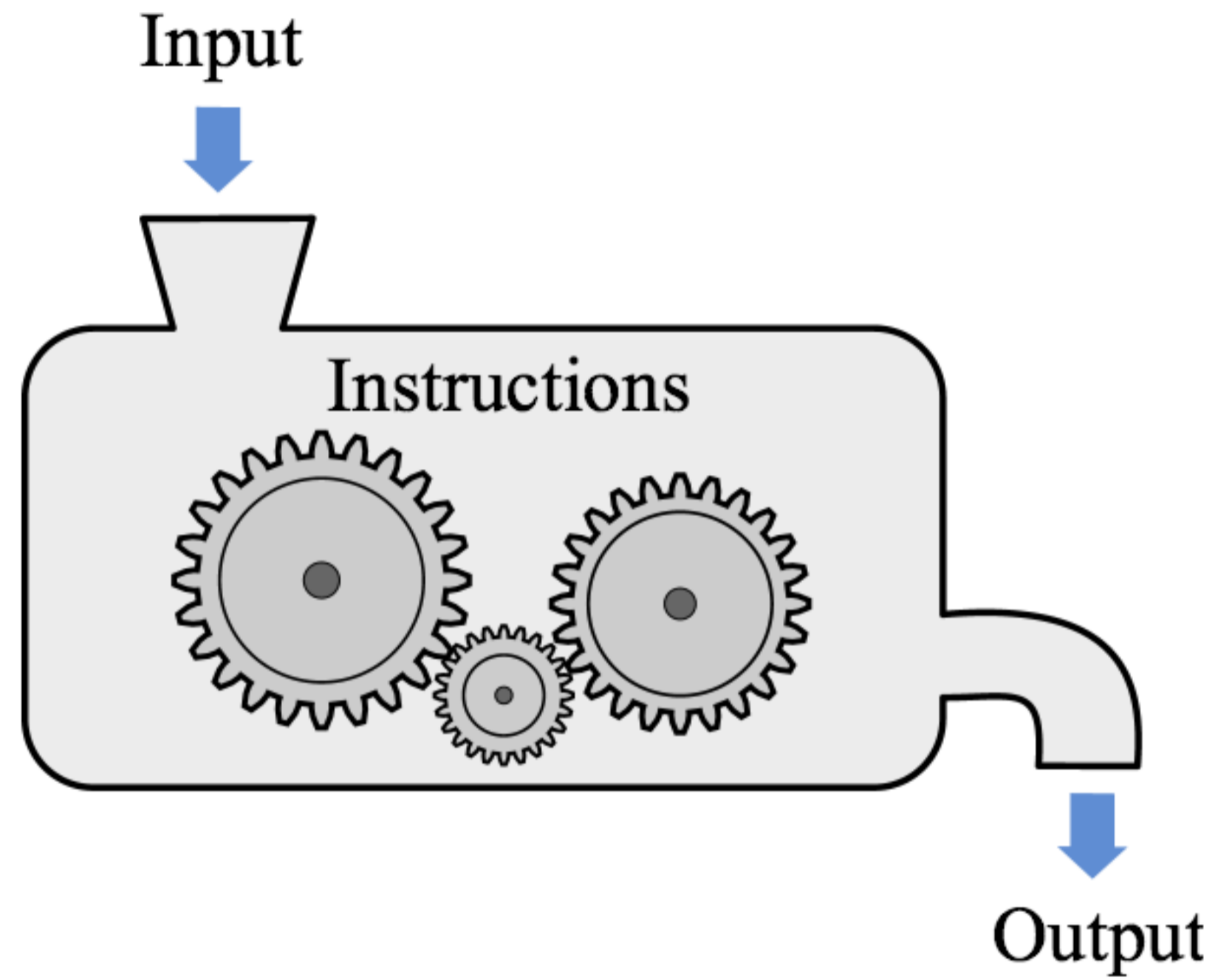
```
col_names = ["colA", "colB", "colC"]
row_names = ["rowD", "rowE", "rowF"]

for col, col_name in enumerate(col_names):
    for row, row_name in enumerate(row_names):
        df_example.iloc[row, col] = col_name+row_name
print(df_example)
```

	colA	colB	colC
rowD	colArowD	colBrowD	colCrowD
rowE	colArowE	colBrowE	colCrowE
rowF	colArowF	colBrowF	colCrowF

# Funksjoner

---



# Repetisjon

---

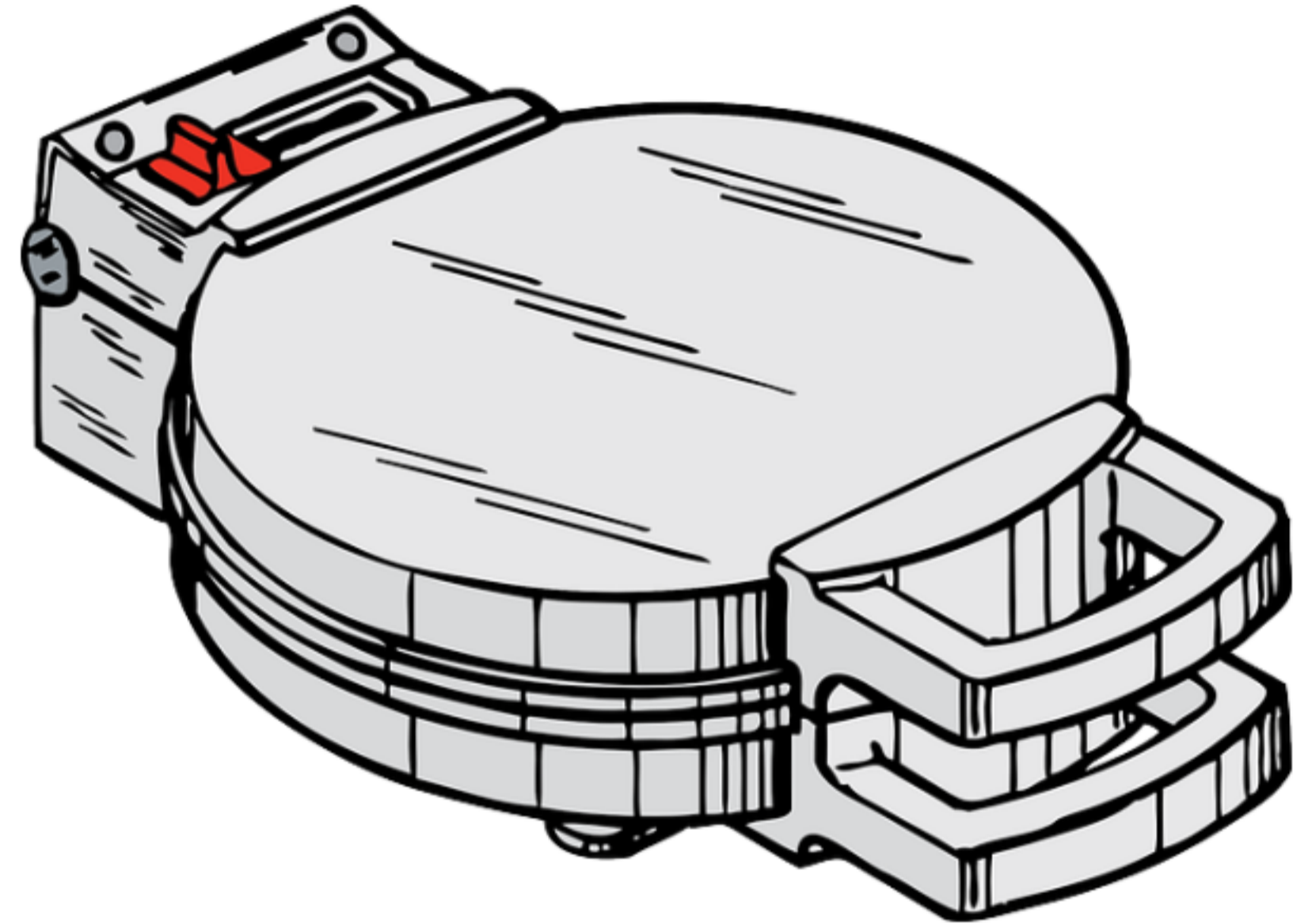
## **Rule of three.**

In programming, there is a rule of thumb called "the rule of three". It states that if a task has to be repeated three times or more, we should automate it.

# Funksjoner

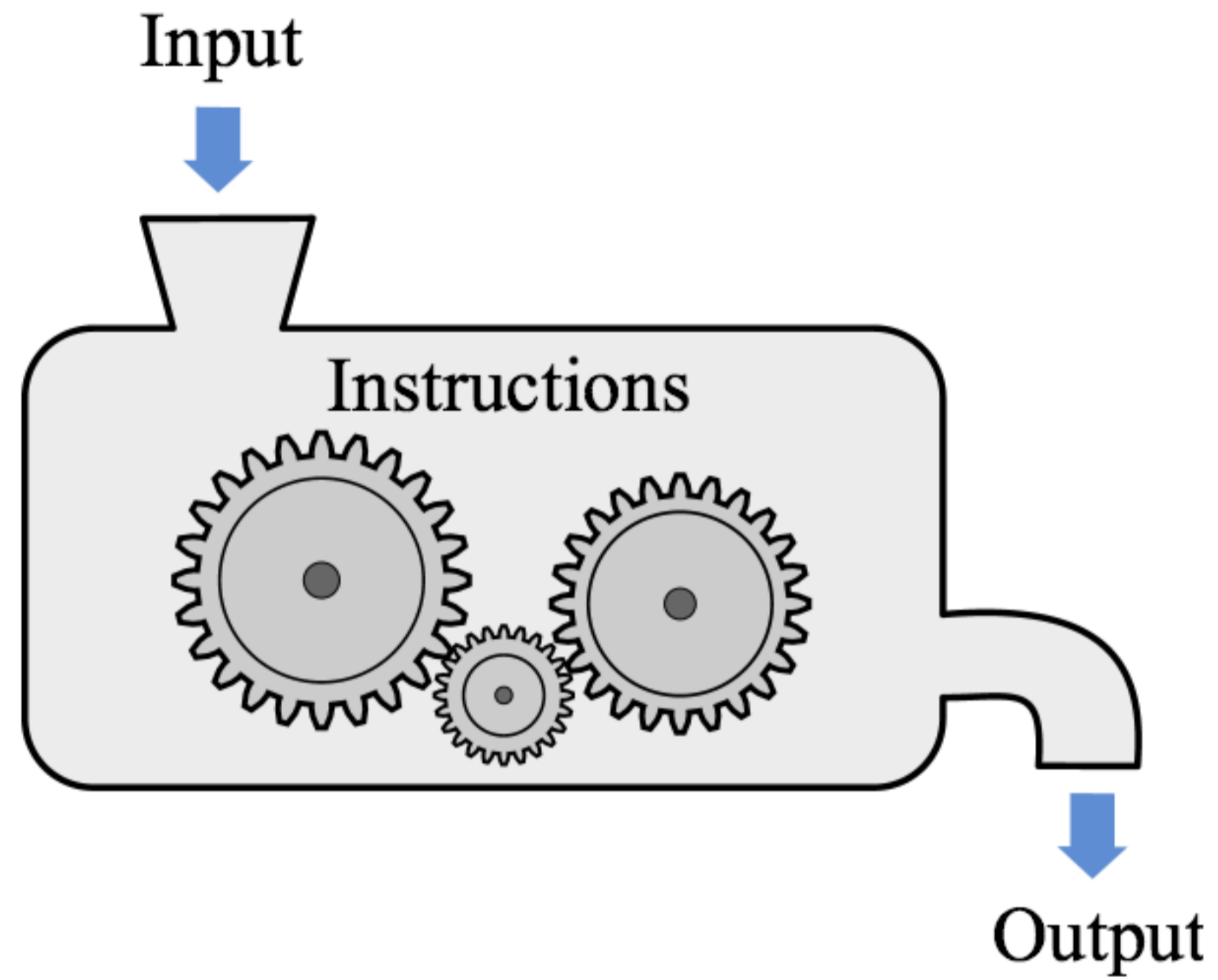
---

Hva er funksjonen til et waffeljern?



# Funksjoner

---



# Funksjoner

---

```
celsius = (fahrenheit - 32)/1.8
```



# Funksjoner

---

```
celsius = (fahrenheit - 32)/1.8
```

```
def fahrenheit_to_celsius(fahrenheit):  
    celsius = (fahrenheit - 32)/1.8  
    return celsius
```

# Funksjoner

---

```
def fahrenheit_to_celsius(fahrenheit):  
    celsius = (fahrenheit - 32)/1.8  
    return celsius
```

```
print(fahrenheit_to_celsius(10))
```

```
-12.222222222222221
```

# Funksjoner

---

```
def fahrenheit_to_celsius(fahrenheit):  
    celsius = (fahrenheit - 32)/1.8  
    return celsius
```

```
T_fahrenheit = 10  
T_celsius = fahrenheit_to_celsius(T_fahrenheit)  
print(T_fahrenheit, "degrees Fahrenheit is", T_celsius, " degrees celsius.")
```

```
10 degrees Fahrenheit is -12.2222222222222221 degrees celsius.
```

# Funksjoner

---

```
def fahrenheit_to_celsius(fahrenheit):  
    celsius = (fahrenheit - 32)/1.8  
    return celsius  
  
bananas = 10  
volkswagen = fahrenheit_to_celsius(bananas)  
print(bananas, "degrees Fahrenheit is", volkswagen, " degrees celsius.")
```

# Funksjoner

---

```
def hello(darwin):  
    dog = (darwin - 32)/1.8  
    return dog  
  
bananas = 10  
volkswagen = hello(bananas)  
print(bananas, "degrees Fahrenheit is", volkswagen, " degrees celsius.")
```

# Funksjoner

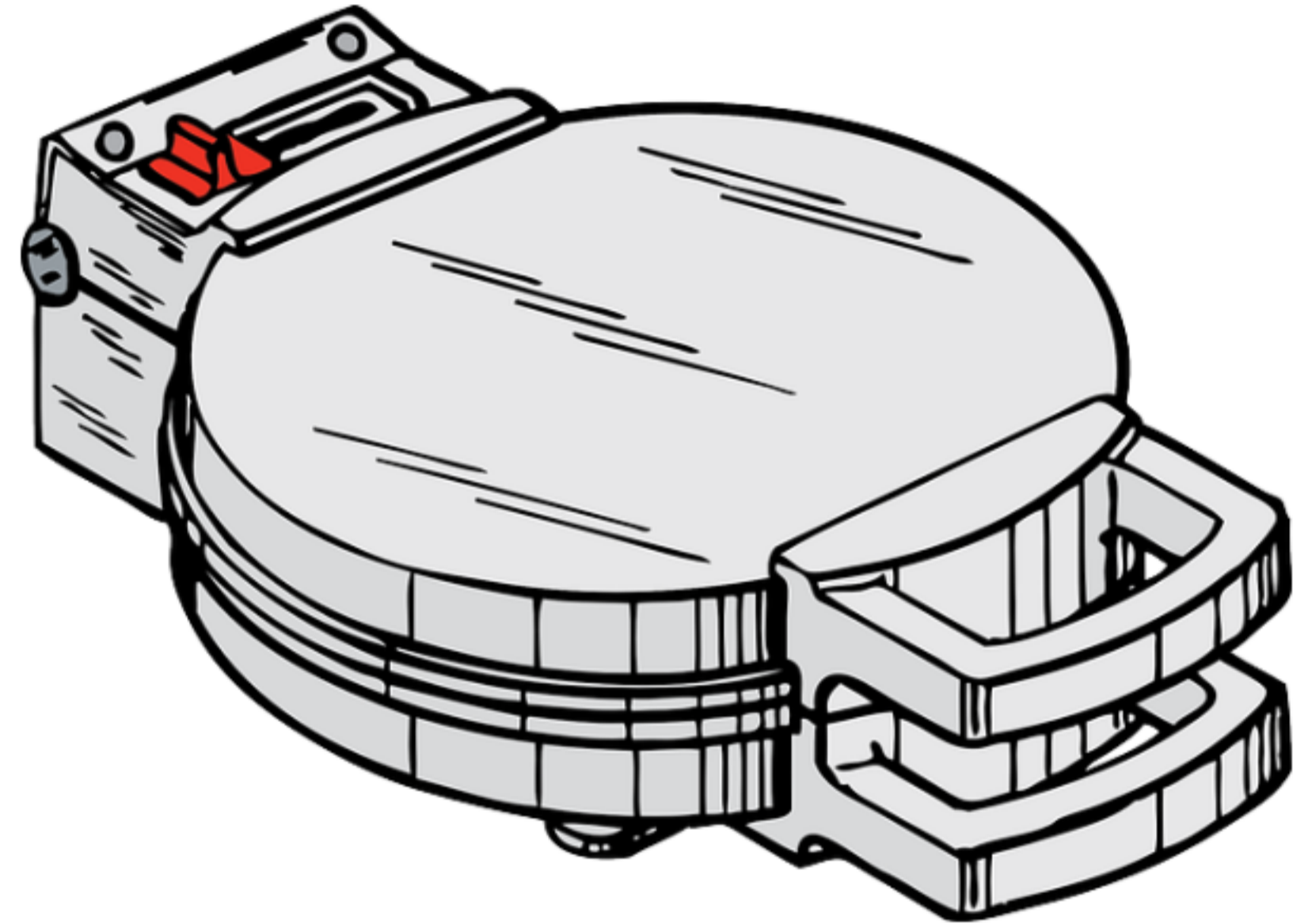
---

```
def fahrenheit_to_celsius(fahrenheit):  
    celsius = (fahrenheit - 32)/1.8  
    return celsius  
  
fahrenheit = 10  
celsius = fahrenheit_to_celsius(fahrenheit)  
print(fahrenheit, "degrees Fahrenheit is", celsius, " degrees celsius.")
```

# Funksjoner

---

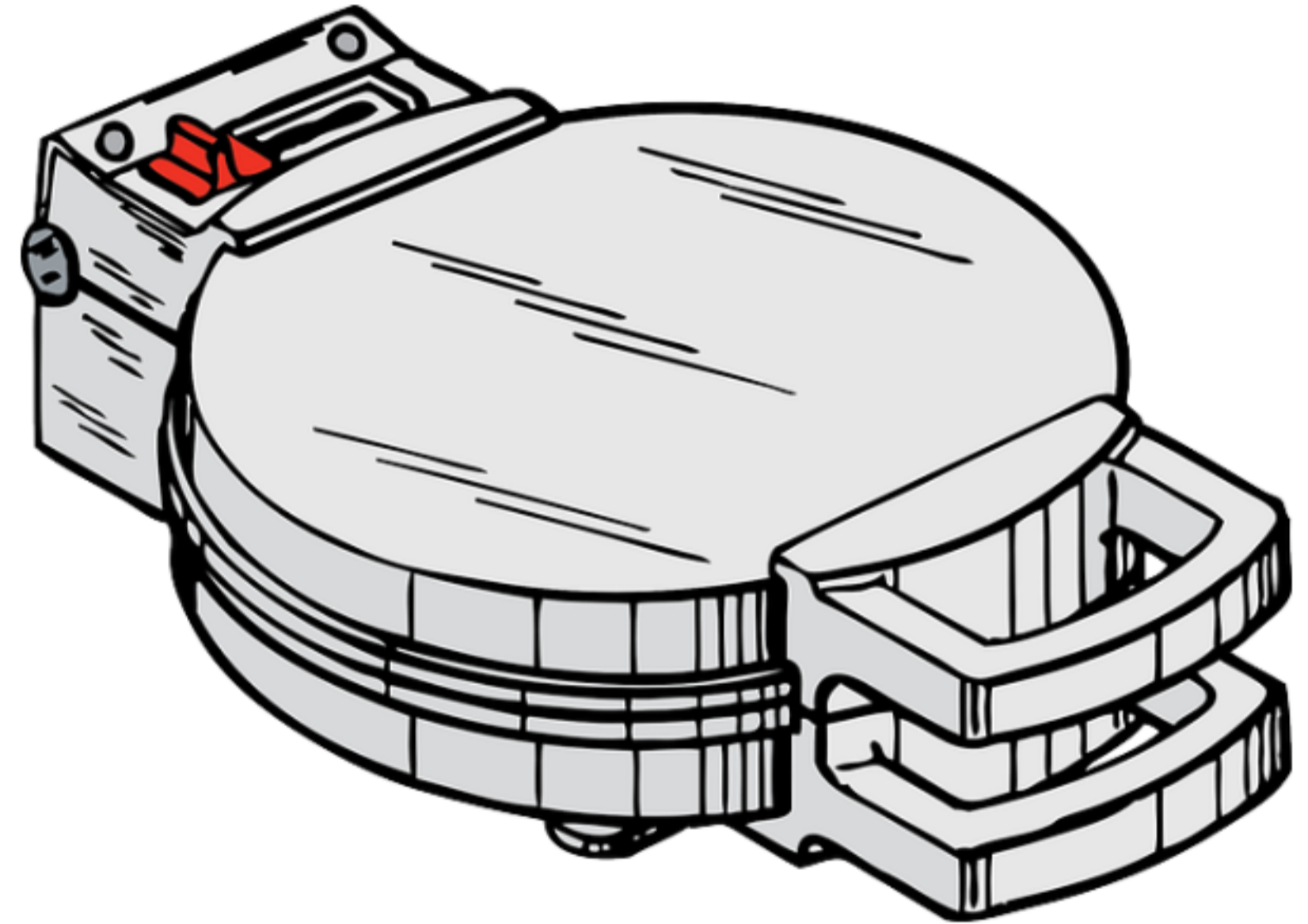
Hva er funksjonen til et waffeljern?



# Funksjoner

---

Hva er funksjonen til et waffeljern?



Waffeljernet bryr seg ikke om deien!



## Punnett square funksjonen

---

```
col_names = ["colA", "colB", "colC"]
row_names = ["rowD", "rowE", "rowF"]

for col, col_name in enumerate(col_names):
    for row, row_name in enumerate(row_names):
        df_example.iloc[row, col] = col_name+row_name
print(df_example)
```

```
def create_punnett_square(parent_1, parent_2):
    punnett_square = pandas.DataFrame(index=parent_1, columns=parent_2)

    for index_1, allele_1 in enumerate(parent_1):
        for index_2, allele_2 in enumerate(parent_2):
            punnett_square.iloc[index_1, index_2] = allele_1 + allele_2

    return punnett_square
```

# Punnett square funksjonen

---

```
parent_1 = ["IG", "ig"]  
parent_2 = ["IG", "ig"]  
  
punnett_square = create_punnett_square(parent_1, parent_2)  
print(punnett_square)
```

```
      IG      ig  
IG  IGIG  IGig  
ig  igIG  igig
```

# Punnett square funksjonen

---

```
parent_1 = ["IG", "Ig", "iG", "ig"]
```

```
parent_2 = ["IG", "Ig", "iG", "ig"]
```

```
punnett_square = create_punnett_square(parent_1, parent_2)
```

```
print(punnett_square)
```

	IG	Ig	iG	ig
IG	IGIG	IGIg	IGiG	IGig
Ig	IgIG	IgIg	IgiG	Igig
iG	iGIG	iGIg	iGiG	iGig
ig	igIG	igIg	igiG	igig

# Mendel's rules

---

## Mendel's rules for inheritance

- An inheritable trait is carried by a gene, and the genes exist in two alleles.
- Each organism inherits two alleles, one from each parent.
- Inherited alleles are chosen at random.
- If the inherited alleles are different, one is dominant and overrules the other, which is recessive.

# Mendel's rules

---

## Mendel's rules for inheritance

- An inheritable trait is carried by a gene, and the genes exist in two alleles.

```
parent_1 = ["B", "b"]
```

# Mendel's rules

---

## Mendel's rules for inheritance

- Each organism inherits two alleles, one from each parent.

```
parent_1 = ["B", "b"]  
parent_2 = ["B", "b"]
```

# Mendel's rules

---

## Mendel's rules for inheritance

- Inherited alleles are chosen at random.

## Tilfeldig valg

---

```
from pylab import *  
  
colors = ["red", "green", "blue", "violet"]  
color = choice(colors)  
  
print(color)
```



# Mendel modellen

---

```
parent_1 = ["B", "b"]
parent_2 = ["B", "b"]

allele_1 = choice(parent_1)
allele_2 = choice(parent_2)

genotype = [allele_1, allele_2]

print("The genotype is:", genotype)
```

# Mendel's rules

---

## Mendel's rules for inheritance

- Inherited alleles are chosen at random.

```
allele_1 = choice(parent_1)
allele_2 = choice(parent_2)
```

# Mendel's rules

---

## Mendel's rules for inheritance

- If the inherited alleles are different, one is dominant and overrules the other, which is recessive.

# Mendel modellen

---

```
parent_1 = ["B", "b"]
parent_2 = ["B", "b"]

allele_1 = choice(parent_1)
allele_2 = choice(parent_2)

genotype = [allele_1, allele_2]

print("The genotype is:", genotype)
```

# Mendel modellen

---

```
parent_1 = ["B", "b"]
parent_2 = ["B", "b"]

allele_1 = choice(parent_1)
allele_2 = choice(parent_2)

genotype = [allele_1, allele_2]

print("The genotype is:", genotype)

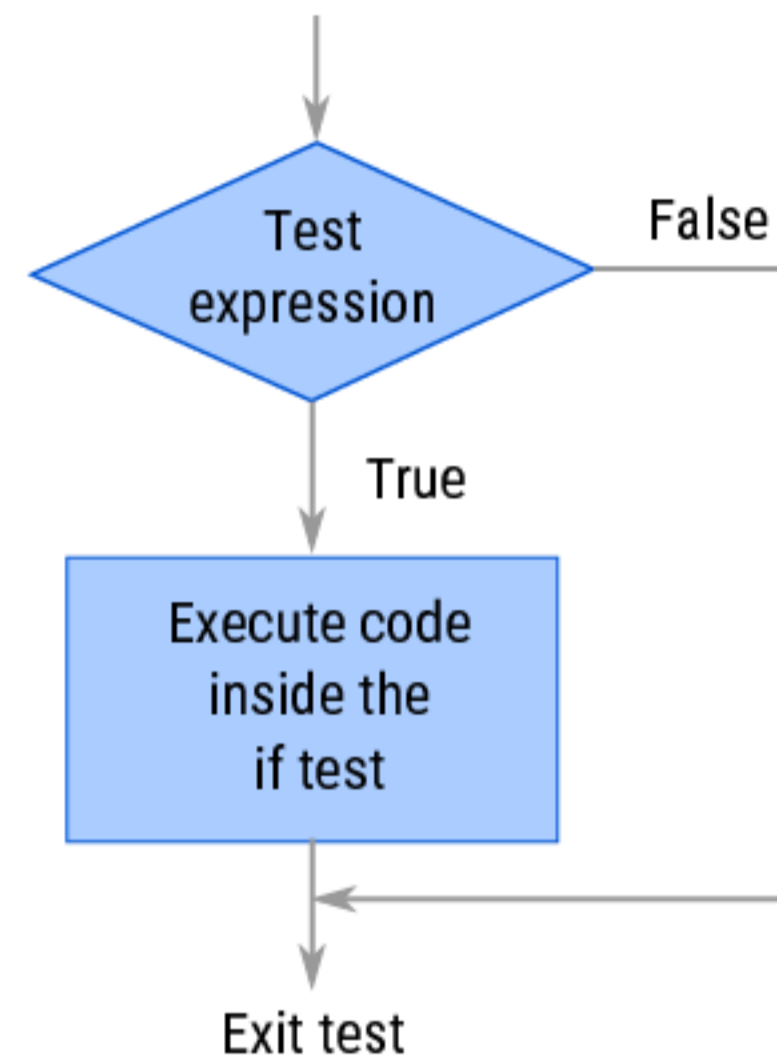
if genotype == ["B", "B"]:
    phenotype = "violet"
if genotype == ["B", "b"]:
    phenotype = "violet"
if genotype == ["b", "B"]:
    phenotype = "violet"
if genotype == ["b", "b"]:
    phenotype = "white"

print("The phenotype is", phenotype)
```

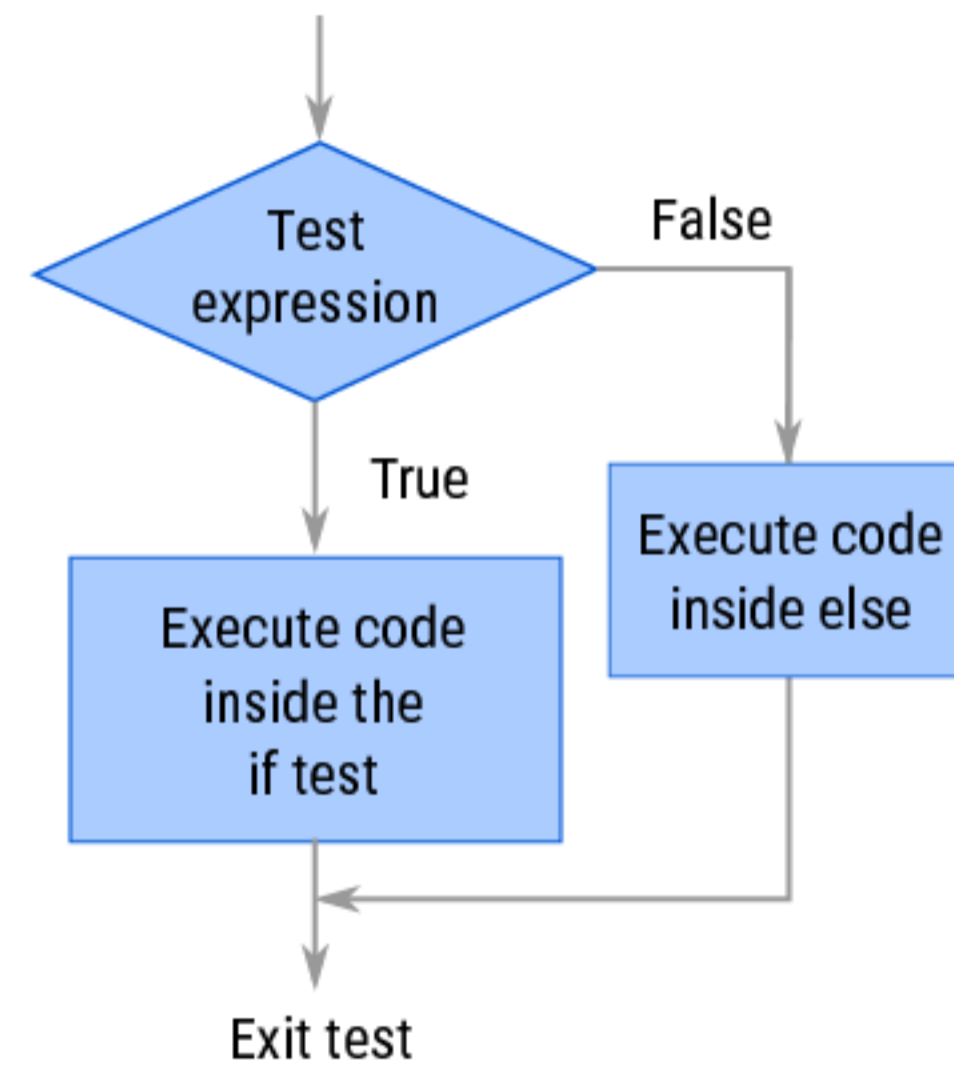
# if .. else

---

## If test



## If-else test



# Mendel modellen

---

```
parent_1 = ["B", "b"]
parent_2 = ["B", "b"]

allele_1 = choice(parent_1)
allele_2 = choice(parent_2)

genotype = [allele_1, allele_2]

print("The genotype is:", genotype)

if genotype == ["b", "b"]:
    phenotype = "white"
else:
    phenotype = "violet"

print("The phenotype is", phenotype)
```

## Gjøre valg - if

---

```
attendees = ['Ola', 'Kari', 'Jane', 'John']  
  
if 'Kari' in attendees:  
    print("Kari is coming!")
```



# Mendel modellen

---

```
genotype = ["b", "B"]

if "B" in genotype:
    phenotype = "violet"
else:
    phenotype = "white"

print("The phenotype is", phenotype)
```

```
The phenotype is violet
```

# Mendel modellen

---

```
genotype = ["b", "b"]

if "B" in genotype:
    phenotype = "violet"
else:
    phenotype = "white"

print("The phenotype is", phenotype)
```

```
The phenotype is white
```

# Mendel's rules

---

## Mendel's rules for inheritance

- If the inherited alleles are different, one is dominant and overrules the other, which is recessive.

```
if "B" in genotype:  
    phenotype = "violet"  
else:  
    phenotype = "white"
```

# Mendel modellen

---

```
parent_1 = ["B", "b"]
parent_2 = ["B", "b"]

allele_1 = choice(parent_1)
allele_2 = choice(parent_2)

genotype = [allele_1, allele_2]

print("The genotype is:", genotype)

if "B" in genotype:
    phenotype = "violet"
else:
    phenotype = "white"

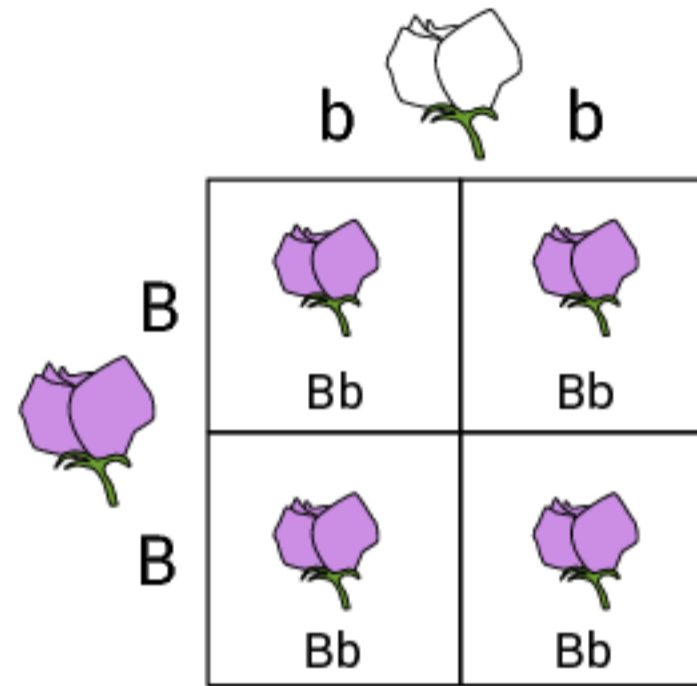
print("The phenotype is", phenotype)
```

# Mendel modellen

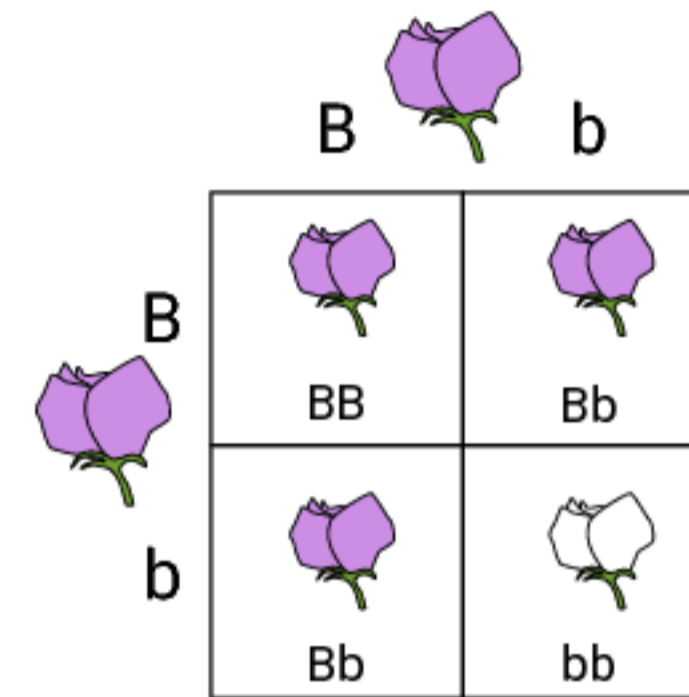
P generation



F<sub>1</sub> generation



F<sub>2</sub> generation



# Mendel modellen

---

Funksjon `create_offspring`

```
def create_offspring(parent_1, parent_2):  
    allele_1 = choice(parent_1)  
    allele_2 = choice(parent_2)  
  
    genotype = [allele_1, allele_2]  
  
    return genotype
```

# Mendel modellen

---

```
violet_flowers_count = 0
white_flowers_count = 0

P_generation_violet = ['B', 'B']
P_generation_white = ['b', 'b']

# Cross-pollinate
F1_generation = create_offspring(P_generation_violet, P_generation_white)

# Self-pollinate
F2_generation = create_offspring(F1_generation, F1_generation)

# Find the phenotype and increase correct count
if 'B' in F2_generation:
    violet_flowers_count = violet_flowers_count + 1
else:
    white_flowers_count = white_flowers_count + 1
```

# Mendel modellen

```
white_flowers_count = 0
violet_flowers_count = 0

for i in range(28000):
    # True bred flowers
    P_generation_violet = ['B', 'B']
    P_generation_white = ['b', 'b']

    # Cross-pollinate
    F1_generation = create_offspring(P_generation_violet, P_generation_white)

    # Self-pollinate
    F2_generation = create_offspring(F1_generation, F1_generation)

    # Find the phenotype and increase correct count
    if 'B' in F2_generation:
        violet_flowers_count = violet_flowers_count + 1
    else:
        white_flowers_count = white_flowers_count + 1
```



## Mendel modellen

---

```
ratio = violet_flowers_count/white_flowers_count

print("The result for the F2 was", violet_flowers_count,
      "violet flowers and", white_flowers_count, "white flowers.")
print("The ratio is", ratio, "to 1")
```

## Mendel modellen

---

```
ratio = violet_flowers_count/white_flowers_count

print("The result for the F2 was", violet_flowers_count,
      "violet flowers and", white_flowers_count, "white flowers.")
print("The ratio is", ratio, "to 1")
```

```
The result for the F2 was 21014 violet flowers and 6986 white flowers.
The ratio is 3.0080160320641283 to 1
```

## Utvalgte øvelser

---

Bruke Google for å finne svar på spørsmålet:

> why doesn't `range` in python return a list?

## Utvalgte øvelser

---

- Exercise 4: The Rice And Chessboard Story revisited
- Exercise 5: Rabbit population growth